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U.S. APPLICATION NUMBER NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
10/526,533	Shmuel Ben-Sasson	BEN-SASSON13A

INTERNATIONAL APPLICATION NO.
PCT/US03/27332

I.A. FILING DATE	PRIORITY DATE
09/03/2003	09/03/2002

001444
BROWDY AND NEIMARK, P.L.L.C.
624 NINTH STREET, NW
SUITE 300
WASHINGTON, DC 20001-5303

CONFIRMATION NO. 6726
371 FORMALITIES LETTER

OC000000018489451

Date Mailed: 04/10/2006

NOTIFICATION OF DEFECTIVE RESPONSE

The following items have been submitted by the applicant or the IB to the United States Patent and Trademark Office as a Designated / Elected Office (37 CFR 1.495)

DEF.RES.NON.EXT.= 10/MY/2006

- Indication of Small Entity Status
- Priority Document
- Copy of the International Application filed on 03/03/2005
- Copy of the International Search Report filed on 03/03/2005
- Information Disclosure Statements filed on 03/03/2005
- Biochemical Sequence Diskette filed on 01/18/2006
- Oath or Declaration filed on 01/18/2006
- Biochemical Sequence Listing filed on 01/18/2006
- Request for Immediate Examination filed on 03/03/2005
- U.S. Basic National Fees filed on 03/03/2005
- Priority Documents filed on 03/03/2005

Applicant's response filed 01/18/2006 is hereby acknowledged. The following requirements set forth in the NOTIFICATION of MISSING REQUIREMENTS mailed 09/19/2005 have not been completed.

The applicant needs to satisfy supplemental fees problems indicated below.

The following items **MUST** be furnished within the period set forth below in order to complete the requirements for acceptance under 35 U.S.C. 371:

- Additional claim fees of **$455** as a small entity, including any required multiple dependent claim fee, are required. Applicant must submit the additional claim fees or cancel the additional claims for which fees are due.

SUMMARY OF FEES DUE:

Total additional fees required for this application is **$455** for a Small Entity:

- The paper or compact disc copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e). Applicant must provide a substitute paper or compact disc copy of the "Sequence Listing", **as well as an amendment specifically directing its entry into the application** OR a substitute computer readable form (CRF) copy of the "Sequence Listing". These two items must be the same. Applicant must also provide a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d). If the effective filing date is on or after September 8, 2000, see the final rulemaking notice published in the Federal Register at 65 FR 54604 (September 8, 2000) and 1238 OG 145 (September 19, 2000).
- A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

Applicant is required to complete the response within a time limit of ONE MONTH from the date of this Notification or within the time remaining in the response set forth in the Notification of Missing Requirements, whichever is the longer. No extension of this time limit may be granted under 37 CFR 1.136, but the period for response set in the Notification of Missing Requirements may be extended under 37 CFR 1.136(a).

- Total additional claim fee(s) for this application is $ **455**

Applicant is cautioned that correction of the above items may cause the specification and drawings page count to exceed 100 pages. If the specification and drawings exceed 100 pages, applicant will need to submit the required application size fee.

For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:

- **For Rules Interpretation, call (571) 272-0951**
- **For Patentin Software Program Help, call Patent EBC at 1-866-217-9197 or directly at 703-305-3028 / 703-308-6845 between the hours of 6 a.m. and 12 midnight, Monday through Friday, EST.**
- **Send e-mail correspondence for Patentin Software Program Help @ ebc@uspto.gov**

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

A copy of this notice MUST be returned with the response.

CHARITTA A BURT

Telephone: (703) 308-9140 EXT 207

PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
10/526,533	PCT/US03/27332	BEN-SASSON13A

FORM PCT/DO/EO/916 (371 Formalities Notice)

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _10/526,533_

Source: _PCT_

Date Processed by STIC: _03/09/2006_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER
VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
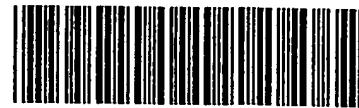Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio **(<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission
   User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
   Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _10/526,533_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics
Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length
The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3____Misaligned Amino
Numbering
The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

4____Non-ASCII
The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5____Variable Length
Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0
"bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7____Skipped Sequences
(OLD RULES)
Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8____Skipped Sequences
(NEW RULES)
Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's
(NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10____Invalid <213>
Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

11____Use of <220>
Sequence(s) ___ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

12____PatentIn 2.0
"bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____Misuse of n/Xaa
"n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

AMC – STIC Systems Branch – 03/02/06

PCT

```
   3 <110> APPLICANT: BEN-SASSON, Shmuel
   4        REUVENI, Hadas
   6 <120> TITLE OF INVENTION: SHORT PEPTIDES FROM THE '2ND LOOP' OF 7 TRANSMEMBRANE
RECEPTOR
   7        WHICH SELECTIVELY MODULATE SIGNAL TRANSDUCTION
   9 <130> FILE REFERENCE: BEN-SASSON13A PCT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/526,533
C--> 11 <141> CURRENT FILING DATE: 2005-03-03
  11 <150> PRIOR APPLICATION NUMBER: 60/407,290
  12 <151> PRIOR FILING DATE: 2002-09-03
  14 <160> NUMBER OF SEQ ID NOS: 13
  16 <170> SOFTWARE: PatentIn version 3.3
  18 <210> SEQ ID NO: 1
  19 <211> LENGTH: 9
  20 <212> TYPE: PRT
  21 <213> ORGANISM: Artificial
  23 <220> FEATURE:
  24 <223> OTHER INFORMATION: synthetic
  26 <400> SEQUENCE: 1
  28 Met Arg Pro Tyr Asp Ala Asn Lys Arg
  29 1               5
  32 <210> SEQ ID NO: 2
  33 <211> LENGTH: 9
  34 <212> TYPE: PRT
  35 <213> ORGANISM: Artificial
  37 <220> FEATURE:
  38 <223> OTHER INFORMATION: synthetic
  40 <400> SEQUENCE: 2
  42 Arg Lys Asn Ala Asp Tyr Pro Arg Met
  43 1               5
  46 <210> SEQ ID NO: 3
  47 <211> LENGTH: 19
  48 <212> TYPE: PRT
  49 <213> ORGANISM: EDG3
  51 <400> SEQUENCE: 3
  53 Glu Arg His Leu Thr Met Ile Lys Met Arg Pro Tyr Asp Ala Asn Lys
  54 1               5                   10                  15
  57 Arg His Arg
  61 <210> SEQ ID NO: 4
  62 <211> LENGTH: 9
  63 <212> TYPE: PRT
  64 <213> ORGANISM: R002L103
  66 <400> SEQUENCE: 4
  68 Met Arg Pro Tyr Asp Ala Asn Lys Arg
```

Does Not Comply
Corrected Diskette Needed

Invalid Response. 22137 Responses can be either Artificial, Unknown or Genus Species. Pls see Item #10 on Error Summary Sheet.

```
 69 1               5
 72 <210> SEQ ID NO: 5
 73 <211> LENGTH: 6
 74 <212> TYPE: PRT
 75 <213> ORGANISM: R002L106
 78 <220> FEATURE:
 79 <221> NAME/KEY: misc_feature
 80 <222> LOCATION: (1)..(1)
 81 <223> OTHER INFORMATION: Xaa is Nle
 83 <400> SEQUENCE: 5
```
W--> `85 Xaa Arg Pro Tyr Asn Ala`
```
 86 1               5
 89 <210> SEQ ID NO: 6
 90 <211> LENGTH: 20
 91 <212> TYPE: PRT
 92 <213> ORGANISM: Lop 2 b3-Adr
 94 <400> SEQUENCE: 6
 96 Asp Arg Tyr Leu Ala Val Thr Asn Pro Leu Arg Tyr Gly Ala Leu Val
 97 1               5                   10                  15
100 Thr Lys Arg Cys
101                20
104 <210> SEQ ID NO: 7
105 <211> LENGTH: 11
106 <212> TYPE: PRT
107 <213> ORGANISM: R013L101
110 <220> FEATURE:
111 <221> NAME/KEY: misc_feature
112 <222> LOCATION: (1)..(1)
113 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
115 <400> SEQUENCE: 7
117 Gly Asn Pro Leu Arg Tyr Gly Ala Leu Val Thr
118 1               5                   10
121 <210> SEQ ID NO: 8
122 <211> LENGTH: 10
123 <212> TYPE: PRT
124 <213> ORGANISM: R013L102
127 <220> FEATURE:
128 <221> NAME/KEY: misc_feature
129 <222> LOCATION: (1)..(1)
130 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
132 <400> SEQUENCE: 8
134 Gly Leu Arg Tyr Gly Ala Leu Val Thr Lys
135 1               5                   10
138 <210> SEQ ID NO: 9
139 <211> LENGTH: 10
140 <212> TYPE: PRT
141 <213> ORGANISM: R013L103
144 <220> FEATURE:
145 <221> NAME/KEY: misc_feature
```

Same Error

```
146 <222> LOCATION: (1)..(1)
147 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
149 <400> SEQUENCE: 9
151 Gly Pro Leu Arg Tyr Gly Ala Leu Val Thr
152 1            5               10
155 <210> SEQ ID NO: 10
156 <211> LENGTH: 9
157 <212> TYPE: PRT
158 <213> ORGANISM: R001L115
161 <220> FEATURE:
162 <221> NAME/KEY: misc_feature
163 <222> LOCATION: (1)..(1)
164 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
166 <400> SEQUENCE: 10
168 Gly Leu Arg Tyr His Ser Ile Val Thr
169 1            5
172 <210> SEQ ID NO: 11
173 <211> LENGTH: 10
174 <212> TYPE: PRT
175 <213> ORGANISM: R001L116
178 <220> FEATURE:
179 <221> NAME/KEY: misc_feature
180 <222> LOCATION: (1)..(1)
181 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
183 <400> SEQUENCE: 11
185 Gly Leu Arg Tyr His Ser Ile Val Lys Thr
186 1            5               10
189 <210> SEQ ID NO: 12
190 <211> LENGTH: 7
191 <212> TYPE: PRT
192 <213> ORGANISM: K024H107
195 <220> FEATURE:
196 <221> NAME/KEY: misc_feature
197 <222> LOCATION: (1)..(1)
198 <223> OTHER INFORMATION: Gly is modified with a myristoyl group.
200 <400> SEQUENCE: 12
202 Gly Leu Leu Arg Arg His Ser
203 1            5
206 <210> SEQ ID NO: 13
207 <211> LENGTH: 9
208 <212> TYPE: PRT
209 <213> ORGANISM: K024H124
211 <400> SEQUENCE: 13
213 Leu Gly Leu Leu Arg Arg His Ser Ile
214 1            5
```

Same Error

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 03/09/2006
PATENT APPLICATION: US/10/526,533   TIME: 12:29:17

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\03092006\J526533.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 1

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2

VERIFICATION SUMMARY   DATE: 03/09/2006
PATENT APPLICATION: US/10/526,533   TIME: 12:29:17

Input Set : **A:\sequence listing.txt**
Output Set: **N:\CRF4\03092006\J526533.raw**

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:85 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0